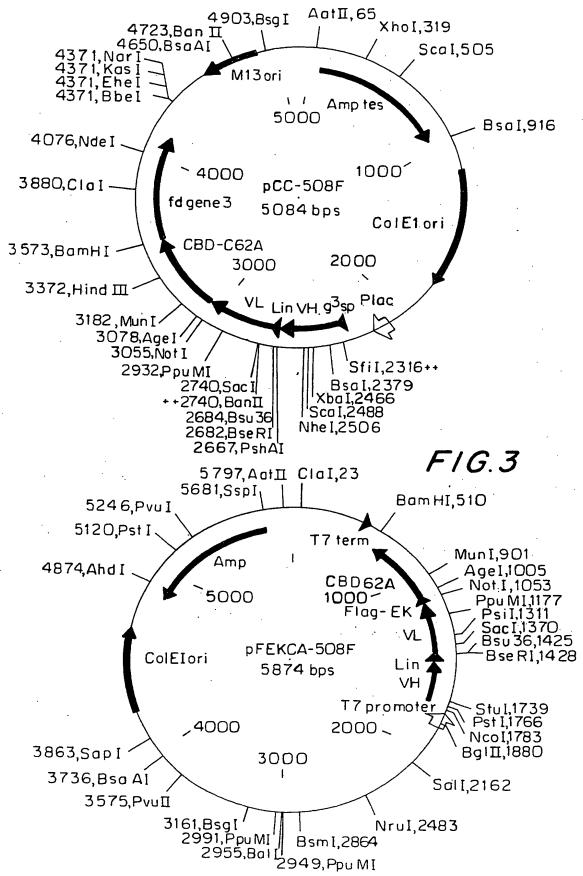


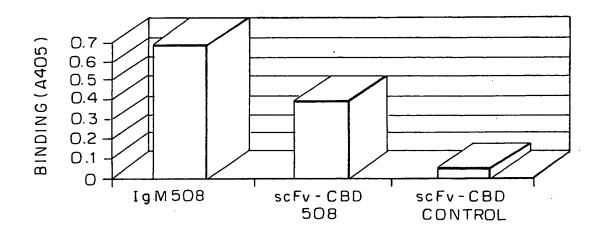


FIG.2

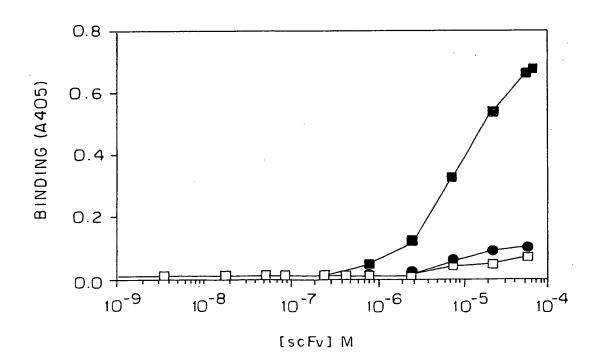




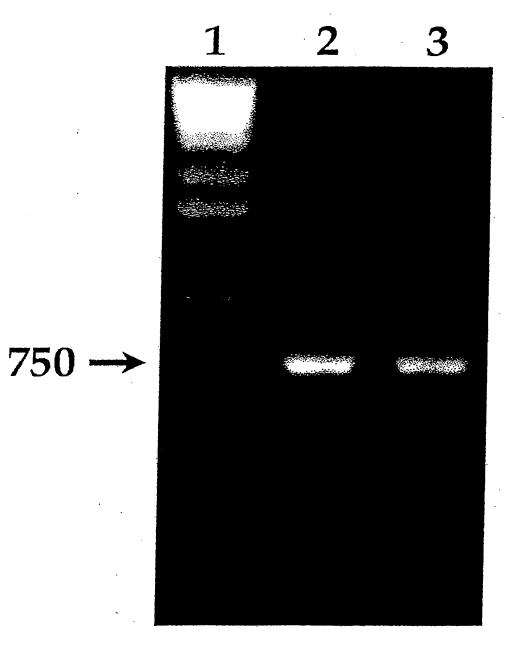
F1G.4



F/G.7



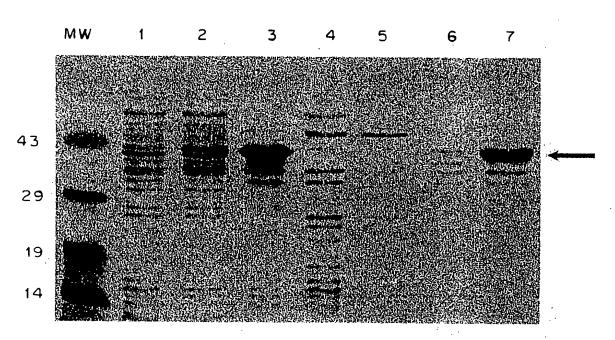




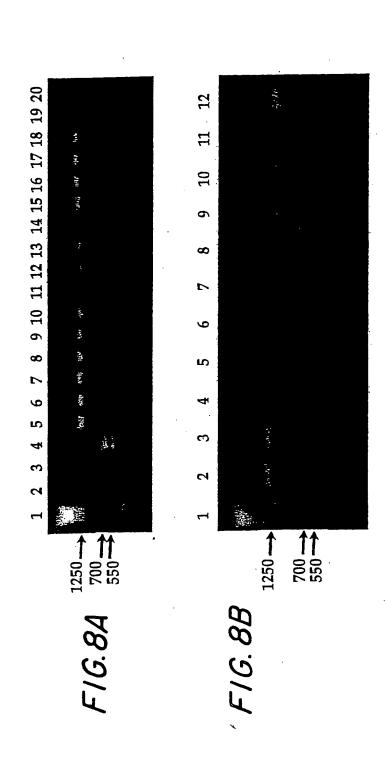
F1G. 5



F1G.6

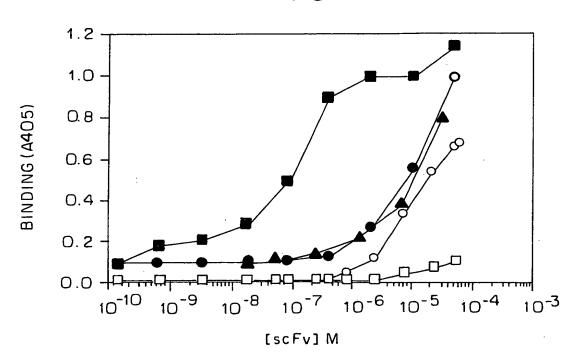




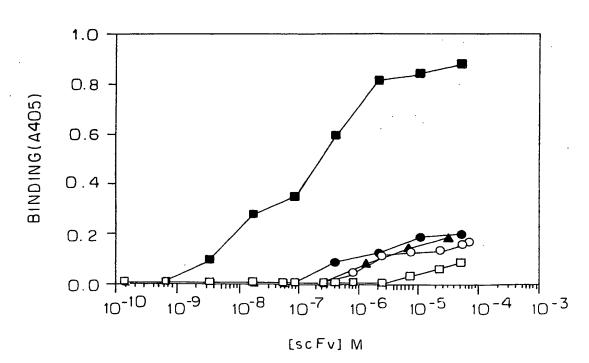




F1G.9a

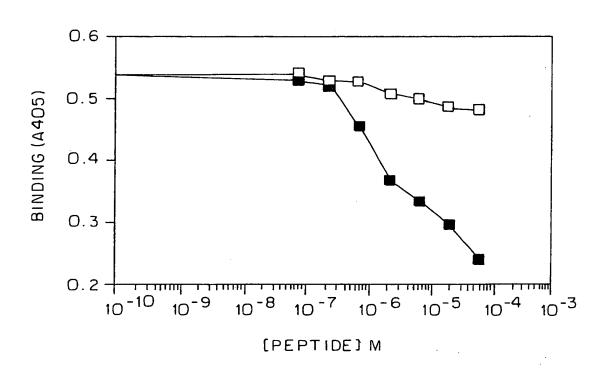








F1G.10



F16.110

CTG GTG AGG CCT GGG GTC TCA GTG AAG AIT ile ser val CAG GTC AAA CTG CAG GAG TCA GGG GCT GAG CTG GTG AGG CCT GGG GTC gln val lys leu gln glu ser gly ala glu leu val arg pro gly val

CAG AGT gln ser ACT GAT TAT GCT ATG CAC TGG GTG AAG thr asp tyr ala met his trp val lys GGC TAC ACA TTC gly tyr thr phe GGT TCT ser cys lys gly ser AAG TGC TCC

GGA GTT ATT AGT ACT TAC TAT GGT GAT GCT AGC TAC gly val ile ser thr tyr tyr gly asp ala ser tyr CDR 1 GAG TGG ATT glu trp ile GCA AAG AGT CTA ala lys ser leu CAT his

AAC CAG AAG TTC AAG GGC AAG GCC ACA ATG ACT GTA GAC AAA TCC TCC AGC ACA GCC TAT asn gln lys phe lys gly lys ala thr met thr val asp lys ser ser thr ala tyr CDR 2 CDR 2

glu leu ala arg leu thr ser glu asp ser ala ile tyr tyr cys ala arg <u>gly ala</u> GAA CTT GCC AGA CTG ACA TCT GAG GAT TCT GCC ATC TAT TAC TGT GCA AGA ATG met

ACT ATG TCC TAC TTT GAC TAC TGG GGC CAA GTG ACC ACG GTC ACC GTC TCC TCA ggt gga thr met ser tyr phe asp tyr trp gly gln val thr thr val thr val ser ser gly gly CDR 3 CDR 3

F1G.11b

ggc ggt tca ggc gga gtt ggc tct ggc ggt ggc gga tcg GAC ATC GAG CTC ACT CAG TCT gly gly gly ser gly gly gly ser asp ile glu leu thr gln ser

Linker

CCA GCA ATC ATG TCT GCA TCT CCA GGG GAG AAG GTC ACC ATG ACC TGC AGT GCC AGC TCA pro ala ile met ser ala ser pro gly glu lys val thr met thr cys <u>ser ala ser ser</u>

pro lys arg trp ile AAA AGA TGG ATT CCC TCC ser AGT ATA AGT TAC ATG CAC TGG TAT CAG CAG AAG CCA GGC ACC ser ile ser tyr met his trp tyr gln gln lys pro gly thr CDR 1

GGA GTC CCT GCT CGC TTC AGT GGC AGT GGG TCT GGG gly val pro ala arg phe ser gly ser gly ser gly asp thr ser lys leu ala ser TAT GAC ACA TCC AAA CTG GCT TCT tyr

GAG GCT GAA GAT GCT GCC ACT TAT TAC TGC glu ala glu asp ala ala thr tyr tyr cys ACC TCT TAT TCT CTC ACA ATC AGC AGC ATG thr ser tyr ser leu thr ile ser ser met CDR 2

CAT CAG CCG AGT AGT TAC CCA TTC ACG TTC GGA GGG GGG GCC AAG CTG GAA ATA AAA his qln arg ser ser tyr pro phe thr phe gly gly gly ala lys leu glu ile lys CDR 3



F I G. 12

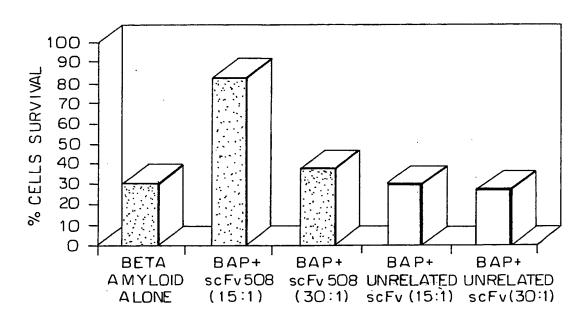
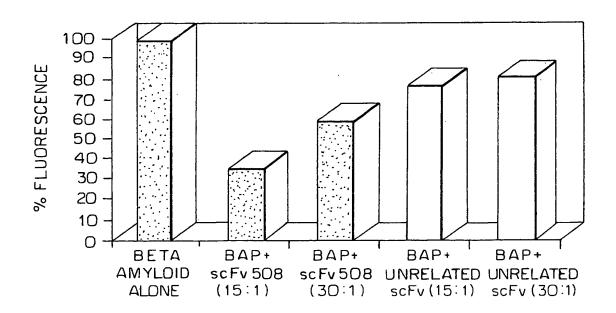


FIG. 13





F1G. 140

F16.14c



F16.14d



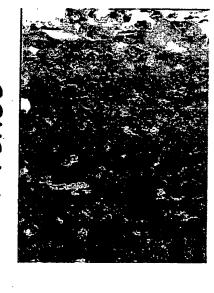




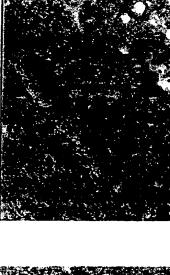
F16.154

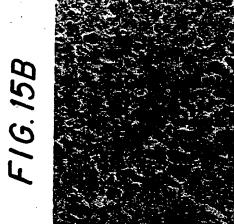


F16.15C

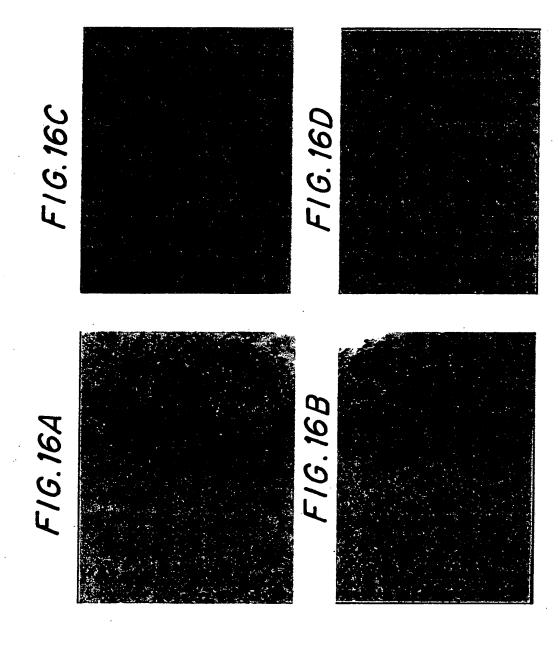


F16.15D

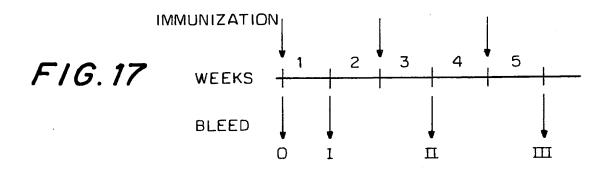


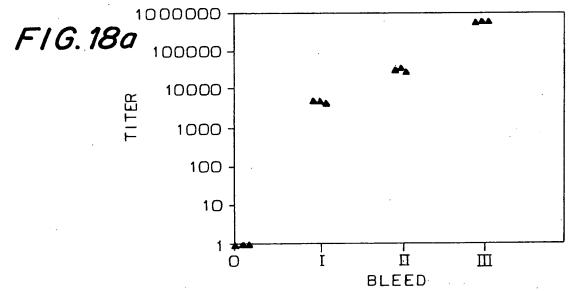


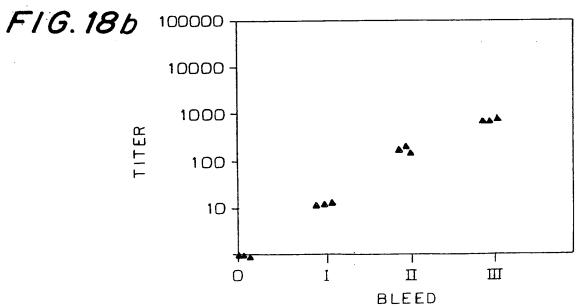






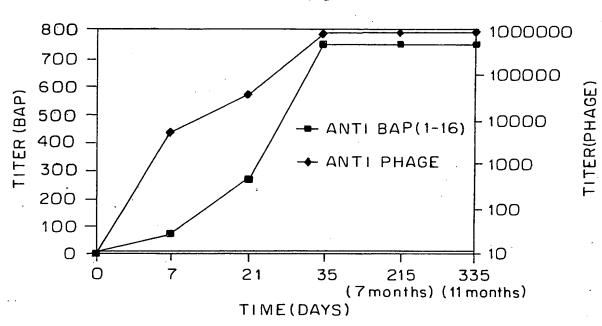




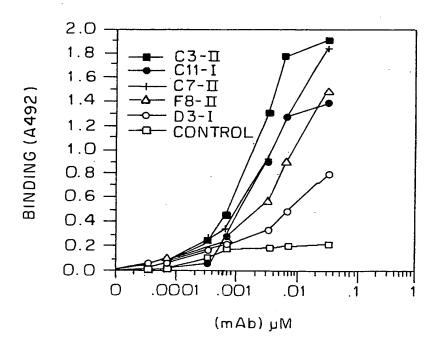




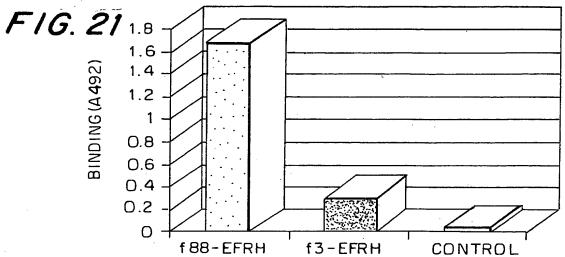
F1G.19



F1G.20



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F1G.22a

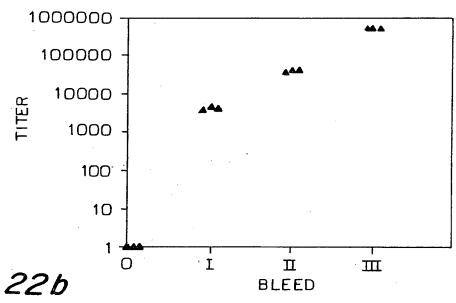
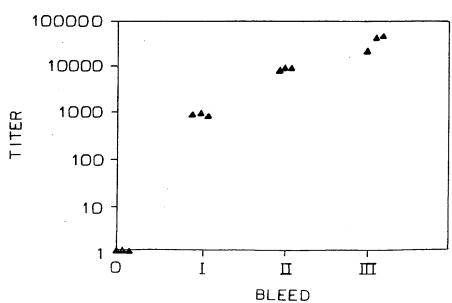
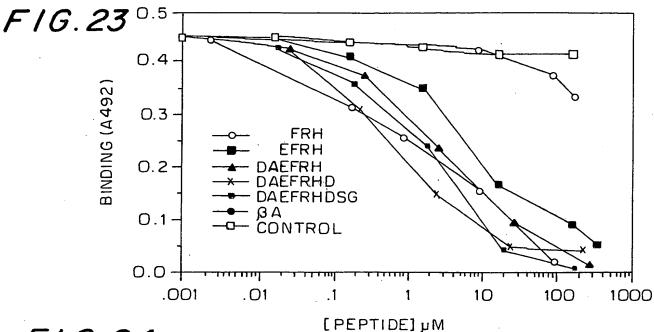


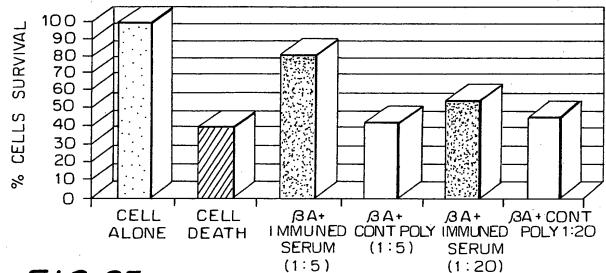
FIG. 22b



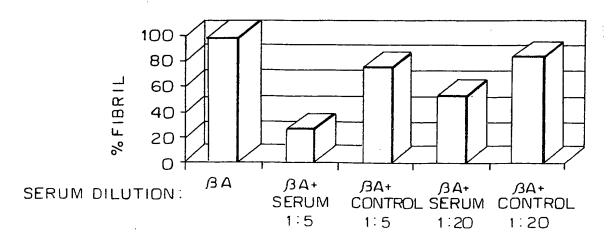




F1G.24



F1G.25

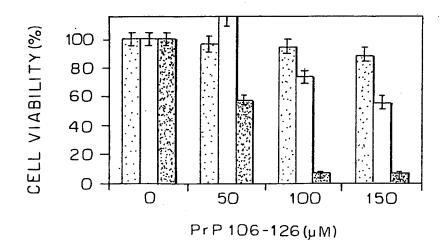




F1G.26

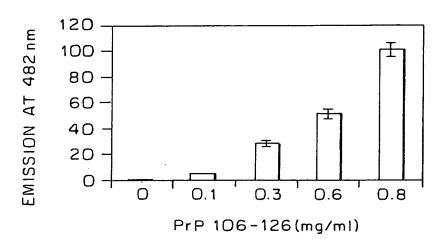
HUMAN Prp 106-126: KTNMKHMAGAAAAGAVVGGLG MOUSE Prp 105-125: KTNLKHVAGAAAAGAVVGGLG

F1G.27

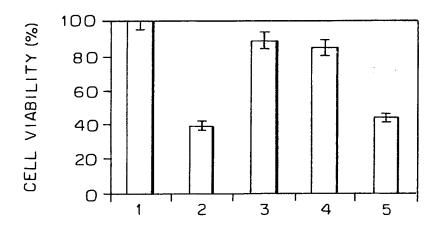




F1G.28

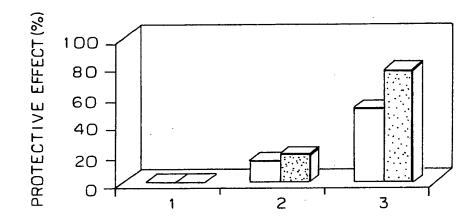


F1G.29





F1G.30



F1G.31

